

## **STIC Biotechnology Systems Branch**

### **RAW SEQUENCE LISTING ERROR REPORT**

**COPY**

**The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:**

Application Serial Number: 10/575,261  
Source: IFWP  
Date Processed by STIC: 04/24/2006

**THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.**

**PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:**

- 1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

**FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221**

**TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:**

**<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>**

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

COPY



IFWP

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/575,261

DATE: 04/24/2006  
TIME: 16:15:41

Input Set : A:\seq list.txt  
Output Set: N:\CRF4\04242006\J575261.raw

3 <110> APPLICANT: KYOWA HAKKO KOGYO CO., LTD.  
5 <120> TITLE OF INVENTION: Fusion protein composition  
7 <130> FILE REFERENCE: 11613WO1  
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/575,261  
C--> 9 <141> CURRENT FILING DATE: 2006-04-10  
9 <150> PRIOR APPLICATION NUMBER: P2003-350158  
10 <151> PRIOR FILING DATE: 2003-10-08  
12 <160> NUMBER OF SEQ ID NOS: 113  
14 <170> SOFTWARE: PatentIn Ver. 2.1

## ERRORED SEQUENCES

131 <210> SEQ ID NO: 2 → found 372  
 132 <211> LENGTH: 376  
 133 <212> TYPE: PRT  
 134 <213> ORGANISM: Cricetulus griseus  
 136 <400> SEQUENCE: 2  
 137 Met Ala His Ala Pro Ala Ser Cys Pro Ser Ser Arg Asn Ser Gly Asp  
 138       1                       5                           10                       15  
 140 Gly Asp Lys Gly Lys Pro Arg Lys Val Ala Leu Ile Thr Gly Ile Thr  
 141       20                      25                           30  
 143 Gly Gln Asp Gly Ser Tyr Leu Ala Glu Phe Leu Leu Glu Lys Gly Tyr  
 144       35                      40                           45  
 146 Glu Val His Gly Ile Val Arg Arg Ser Ser Ser Phe Asn Thr Gly Arg  
 147       50                      55                           60  
 149 Ile Glu His Leu Tyr Lys Asn Pro Gln Ala His Ile Glu Gly Asn Met  
 150       65                      70                           75                       80  
 152 Lys Leu His Tyr Gly Asp Leu Thr Asp Ser Thr Cys Leu Val Lys Ile  
 E--> 153 85                   85 90                   90 95                   95 100  
 155 Ile Asn Glu Val Lys Pro Thr Glu Ile Tyr Asn Leu Gly Ala Gln Ser  
 E--> 156 100 105           105 110           110 115  
 158 His Val Lys Ile Ser Phe Asp Leu Ala Glu Tyr Thr Ala Asp Val Asp  
 E--> 159 115 120           120 125           125 130  
 161 Gly Val Gly Thr Leu Arg Leu Leu Asp Ala Ile Lys Thr Cys Gly Leu  
 E--> 162 130 135           135 140           140 145  
 164 Ile Asn Ser Val Lys Phe Tyr Gln Ala Ser Thr Ser Glu Leu Tyr Gly  
 E--> 165 145 150           150 155           155 160           160  
 167 Lys Val Gln Glu Ile Pro Gln Lys Glu Thr Thr Pro Phe Tyr Pro Arg  
 E--> 168 165 165 170           170 175           175 180  
 170 Ser Pro Tyr Gly Ala Ala Lys Leu Tyr Ala Tyr Trp Ile Val Val Asn  
 E--> 171 180 185           185 190           190 195  
 173 Phe Arg Glu Ala Tyr Asn Leu Phe Ala Val Asn Gly Ile Leu Phe Asn  
 195                           200                           205

Does Not Comply  
Corrected Diskette Needed  
(PG-1,2141516)

Invalid  
Amino Acid  
numbers

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/575,261

DATE: 04/24/2006  
TIME: 16:15:41

**COPY**

Input Set : A:\seq list.txt  
Output Set: N:\CRF4\04242006\J575261.raw

E--> 174	200	205	210
176 His Glu Ser Pro Arg Arg Gly Ala Asn Phe Val Thr Arg Lys Ile Ser			
E--> 177	210 215	215 220	220 225
179 Arg Ser Val Ala Lys Ile Tyr Leu Gly Gln Leu Glu Cys Phe Ser Leu			
E--> 180 225 230	230 235	235 240	240
182 Gly Asn Leu Asp Ala Lys Arg Asp Trp Gly His Ala Lys Asp Tyr Val			
E--> 183 245	245 250	250 255	255 260
185 Glu Ala Met Trp Leu Met Leu Gln Asn Asp Glu Pro Glu Asp Phe Val			
E--> 186	260 265	265 270	270 275
188 Ile Ala Thr Gly Glu Val His Ser Val Arg Glu Phe Val Glu Lys Ser			
E--> 189	275 280	280 285	285 290
191 Phe Met His Ile Gly Lys Thr Ile Val Trp Glu Gly Lys Asn Glu Asn			
E--> 192	290 295	295 300	300 305
194 Glu Val Gly Arg Cys Lys Glu Thr Gly Lys Ile His Val Thr Val Asp			
E--> 195 305 310	310 315	315 320	320
197 Leu Lys Tyr Tyr Arg Pro Thr Glu Val Asp Phe Leu Gln Gly Asp Cys			
E--> 198 325	325 330	330 335	335 340
200 Ser Lys Ala Gln Gln Lys Leu Asn Trp Lys Pro Arg Val Ala Phe Asp			
E--> 201	340 345	345 350	350 355
204 Glu Leu Val Arg Glu Met Val Gln Ala Asp Val Glu Leu Met Arg Thr			
E--> 205	355 360	360 365	365 370
207 Asn Pro Asn Ala			
E--> 208	370 375		
471 <210> SEQ ID NO: 7			
472 <211> LENGTH: 575			
473 <212> TYPE: PRT			
474 <213> ORGANISM: Cricetulus griseus			
476 <400> SEQUENCE: 7			
477 Met Arg Ala Trp Thr Gly Ser Trp Arg Trp Ile Met Leu Ile Leu Phe			
478 1 5 10 15			
480 Ala Trp Gly Thr Leu Leu Phe Tyr Ile Gly Gly His Leu Val Arg Asp			
481 20 25 30			
483 Asn Asp His Pro Asp His Ser Ser Arg Glu Leu Ser Lys Ile Leu Ala			
484 35 40 45			
486 Lys Leu Glu Arg Leu Lys Gln Gln Asn Glu Asp Leu Arg Arg Met Ala			
487 50 55 60			
489 Glu Ser Leu Arg Ile Pro Glu Gly Pro Ile Asp Gln Gly Thr Ala Thr			
490 65 70 75 80			
492 Gly Arg Val Arg Val Leu Glu Glu Gln Leu Val Lys Ala Lys Glu Gln			
493 85 90 95			
495 Ile Glu Asn Tyr Lys Lys Gln Ala Arg Asn Asp Leu Gly Lys Asp His			
496 100 105 110			
498 Glu Ile Leu Arg Arg Arg Ile Glu Asn Gly Ala Lys Glu Leu Trp Phe			
499 115 120 125			
501 Phe Leu Gln Ser Glu Leu Lys Lys Leu Lys Lys Leu Glu Gly Asn Glu			
502 130 135 140			
505 Leu Gln Arg His Ala Asp Glu Ile Leu Leu Asp Leu Gly His His Glu			
506 145 150 155 160			
508 Arg Ser Ile Met Thr Asp Leu Tyr Tyr Leu Ser Gln Thr Asp Gly Ala			

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/575,261

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Input Set : A:\seq list.txt  
Output Set: N:\CRF4\04242006\J575261.raw

**COPY**

509	165	170	175	
511	Gly Glu Trp Arg Glu Lys Glu Ala Lys Asp Leu Thr Glu Leu Val Gln			
512	180	185	190	
514	Arg Arg Ile Thr Tyr Leu Gln Asn Pro Lys Asp Cys Ser Lys Ala Arg			
515	195	200	205	
517	Lys Leu Val Cys Asn Ile Asn Lys Gly Cys Gly Tyr Gly Cys Gln Leu			
518	210	215	220	
520	His His Val Val Tyr Cys Phe Met Ile Ala Tyr Gly Thr Gln Arg Thr			
521	225	230	235	240
523	Leu Ile Leu Glu Ser Gln Asn Trp Arg Tyr Ala Thr Gly Gly Trp Glu			
524	245	250	255	
526	Thr Val Phe Arg Pro Val Ser Glu Thr Cys Thr Asp Arg Ser Gly Leu			
527	260	265	270	
529	Ser Thr Gly His Trp Ser Gly Glu Val Lys Asp Lys Asn Val Gln Val			
530	275	280	285	
532	Val Glu Leu Pro Ile Val Asp Ser Leu His Pro Arg Pro Pro Tyr Leu			
533	290	295	300	
535	Pro Leu Ala Val Pro Glu Asp Leu Ala Asp Arg Leu Leu Arg Val His			
536	305	310	315	320
538	Gly Asp Pro Ala Val Trp Trp Val Ser Gln Phe Val Lys Tyr Leu Ile			
539	325	330	335	
541	Arg Pro Gln Pro Trp Leu Glu Arg Glu Ile Glu Glu Thr Thr Lys Lys			
542	340	345	350	
544	Leu Gly Phe Lys His Pro Val Ile Gly Val His Val Arg Arg Thr Asp			
545	355	360	365	
547	Lys Val Gly Thr Glu Ala Ala Phe His Pro Ile Glu Glu Tyr Met Val			
548	370	375	380	
550	His Val Glu Glu His Phe Gln Leu Leu Glu Arg Arg Met Lys Val Asp			
551	385	390	395	400
553	Lys Lys Arg Val Tyr Leu Ala Thr Asp Asp Pro Ser Leu Leu Lys Glu			
554	405	410	415	
556	Ala Lys Thr Lys Tyr Ser Asn Tyr Glu Phe Ile Ser Asp Asn Ser Ile			
557	420	425	430	
559	Ser Trp Ser Ala Gly Leu His Asn Arg Tyr Thr Glu Asn Ser Leu Arg			
560	435	440	445	
562	Gly Val Ile Leu Asp Ile His Phe Leu Ser Gln Ala Asp Phe Leu Val			
563	450	455	460	
565	Cys Thr Phe Ser Ser Gln Val Cys Arg Val Ala Tyr Glu Ile Met Gln			
566	465	470	475	480
568	Thr Leu His Pro Asp Ala Ser Ala Asn Phe His Ser Leu Asp Asp Ile			
569	485	490	495	
571	Tyr Tyr Phe Gly Gly Gln Asn Ala His Asn Gln Ile Ala Val Tyr Pro			
572	500	505	510	
574	His Gln Pro Arg Thr Lys Glu Glu Ile Pro Met Glu Pro Gly Asp Ile			
575	515	520	525	
577	Ile Gly Val Ala Gly Asn His Trp Asn Gly Tyr Ser Lys Gly Val Asn			
578	530	535	540	
580	Arg Lys Leu Gly Lys Thr Gly Leu Tyr Pro Ser Tyr Lys Val Arg Glu			
581	545	550	555	560

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/575,261

DATE: 04/24/2006  
TIME: 16:15:41

Input Set : A:\seq list.txt  
Output Set: N:\CRF4\04242006\J575261.raw

**COPY**

583 Lys Ile Glu Thr Val Lys Tyr Pro Thr Tyr Pro Glu Ala Glu Lys  
**E--> 584** 565 570 *S75*

587 <210> SEQ ID NO: 8  
 588 <211> LENGTH: 575  
 589 <212> TYPE: PRT  
 590 <213> ORGANISM: Mus musculus  
 592 <400> SEQUENCE: 8

593 Met Arg Ala Trp Thr Gly Ser Trp Arg Trp Ile Met Leu Ile Leu Phe  
 594 1 5 10 15  
 596 Ala Trp Gly Thr Leu Leu Phe Tyr Ile Gly Gly His Leu Val Arg Asp  
 597 20 25 30  
 599 Asn Asp His Pro Asp His Ser Ser Arg Glu Leu Ser Lys Ile Leu Ala  
 600 35 40 45  
 602 Lys Leu Glu Arg Leu Lys Gln Gln Asn Glu Asp Leu Arg Arg Met Ala  
 604 50 55 60  
 606 Glu Ser Leu Arg Ile Pro Glu Gly Pro Ile Asp Gln Gly Thr Ala Thr  
 607 65 70 75 80  
 609 Gly Arg Val Arg Val Leu Glu Glu Gln Leu Val Lys Ala Lys Glu Gln  
 610 85 90 95  
 612 Ile Glu Asn Tyr Lys Lys Gln Ala Arg Asn Gly Leu Gly Lys Asp His  
 613 100 105 110  
 615 Glu Ile Leu Arg Arg Arg Ile Glu Asn Gly Ala Lys Glu Leu Trp Phe  
 616 115 120 125  
 618 Phe Leu Gln Ser Glu Leu Lys Lys Leu Lys His Leu Glu Gly Asn Glu  
 619 130 135 140  
 621 Leu Gln Arg His Ala Asp Glu Ile Leu Leu Asp Leu Gly His His Glu  
 622 145 150 155 160  
 624 Arg Ser Ile Met Thr Asp Leu Tyr Tyr Leu Ser Gln Thr Asp Gly Ala  
 625 165 170 175  
 627 Gly Asp Trp Arg Glu Lys Glu Ala Lys Asp Leu Thr Glu Leu Val Gln  
 628 180 185 190  
 630 Arg Arg Ile Thr Tyr Leu Gln Asn Pro Lys Asp Cys Ser Lys Ala Arg  
 631 195 200 205  
 633 Lys Leu Val Cys Asn Ile Asn Lys Gly Cys Gly Tyr Gly Cys Gln Leu  
 634 210 215 220  
 636 His His Val Val Tyr Cys Phe Met Ile Ala Tyr Gly Thr Gln Arg Thr  
 637 225 230 235 240  
 639 Leu Ile Leu Glu Ser Gln Asn Trp Arg Tyr Ala Thr Gly Gly Trp Glu  
 640 245 250 255  
 642 Thr Val Phe Arg Pro Val Ser Glu Thr Cys Thr Asp Arg Ser Gly Leu  
 643 260 265 270  
 645 Ser Thr Gly His Trp Ser Gly Glu Val Asn Asp Lys Asn Ile Gln Val  
 646 275 280 285  
 648 Val Glu Leu Pro Ile Val Asp Ser Leu His Pro Arg Pro Pro Tyr Leu  
 649 290 295 300  
 651 Pro Leu Ala Val Pro Glu Asp Leu Ala Asp Arg Leu Leu Arg Val His  
 652 305 310 315 320  
 654 Gly Asp Pro Ala Val Trp Trp Val Ser Gln Phe Val Lys Tyr Leu Ile  
 655 325 330 335

RAW SEQUENCE LISTING  
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Output Set: N:\CRF4\04242006\J575261.raw

**COPY**

657 Arg Pro Gln Pro Trp Leu Glu Lys Glu Ile Glu Glu Ala Thr Lys Lys  
 658 340 345 350  
 660 Leu Gly Phe Lys His Pro Val Ile Gly Val His Val Arg Arg Thr Asp  
 661 355 360 365  
 663 Lys Val Gly Thr Glu Ala Ala Phe His Pro Ile Glu Glu Tyr Met Val  
 664 370 375 380  
 666 His Val Glu Glu His Phe Gln Leu Leu Ala Arg Arg Met Gln Val Asp  
 667 385 390 395 400  
 669 Lys Lys Arg Val Tyr Leu Ala Thr Asp Asp Pro Thr Leu Leu Lys Glu  
 670 405 410 415  
 672 Ala Lys Thr Lys Tyr Ser Asn Tyr Glu Phe Ile Ser Asp Asn Ser Ile  
 673 420 425 430  
 675 Ser Trp Ser Ala Gly Leu His Asn Arg Tyr Thr Glu Asn Ser Leu Arg  
 676 435 440 445  
 678 Gly Val Ile Leu Asp Ile His Phe Leu Ser Gln Ala Asp Phe Leu Val  
 679 450 455 460  
 681 Cys Thr Phe Ser Ser Gln Val Cys Arg Val Ala Tyr Glu Ile Met Gln  
 682 465 470 475 480  
 684 Thr Leu His Pro Asp Ala Ser Ala Asn Phe His Ser Leu Asp Asp Ile  
 685 485 490 495  
 687 Tyr Tyr Phe Gly Gly Gln Asn Ala His Asn Gln Ile Ala Val Tyr Pro  
 688 500 505 510  
 690 His Lys Pro Arg Thr Glu Glu Ile Pro Met Glu Pro Gly Asp Ile  
 691 515 520 525  
 693 Ile Gly Val Ala Gly Asn His Trp Asp Gly Tyr Ser Lys Gly Ile Asn  
 694 530 535 540  
 696 Arg Lys Leu Gly Lys Thr Gly Leu Tyr Pro Ser Tyr Lys Val Arg Glu  
 697 545 550 555 560  
 699 Lys Ile Glu Thr Val Lys Tyr Pro Thr Tyr Pro Glu Ala Glu Lys  
 E--> 700 565 570 575  
 2297 <210> SEQ ID NO: 64  
 2298 <211> LENGTH: 235  
 2299 <212> TYPE: PRT  
 2300 <213> ORGANISM: Homo sapiens  
 2302 <400> SEQUENCE: 64  
 2303 Leu Pro Ala Gln Val Ala Phe Thr Pro Tyr Ala Pro Glu Pro Gly Ser  
 2304 1 5 10 15  
 2306 Thr Cys Arg Leu Arg Glu Tyr Tyr Asp Gln Thr Ala Gln Met Cys Cys  
 2307 20 25 30  
 2309 Ser Lys Cys Ser Pro Gly Gln His Ala Lys Val Phe Cys Thr Lys Thr  
 2310 35 40 45  
 2312 Ser Asp Thr Val Cys Asp Ser Cys Glu Asp Ser Thr Tyr Thr Gln Leu  
 2313 50 55 60  
 2315 Trp Asn Trp Val Pro Glu Cys Leu Ser Cys Gly Ser Arg Cys Ser Ser  
 2316 65 70 75 80  
 2318 Asp Gln Val Glu Thr Gln Ala Cys Thr Arg Glu Gln Asn Arg Ile Cys  
 2319 85 90 95  
 2321 Thr Cys Arg Pro Gly Trp Tyr Cys Ala Leu Ser Lys Gln Glu Gly Cys  
 2322 100 105 110

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/575,261

DATE: 04/24/2006  
TIME: 16:15:41

Input Set : A:\seq list.txt  
Output Set: N:\CRF4\04242006\J575261.raw

COPY

2324 Arg Leu Cys Ala Pro Leu Arg Lys Cys Arg Pro Gly Phe Gly Val Ala  
2325 115 120 125  
2327 Arg Pro Gly Thr Glu Thr Ser Asp Val Val Cys Lys Pro Cys Ala Pro  
2328 130 135 140  
2330 Gly Thr Phe Ser Asn Thr Thr Ser Ser Thr Asp Ile Cys Arg Pro His  
2331 145 150 155 160  
2333 Gln Ile Cys Asn Val Val Ala Ile Pro Gly Asn Ala Ser Met Asp Ala  
2334 165 170 175  
2336 Val Cys Thr Ser Thr Ser Pro Thr Arg Ser Met Ala Pro Gly Ala Val  
2337 180 185 190  
2339 His Leu Pro Gln Pro Val Ser Thr Arg Ser Gln His Thr Gln Pro Thr  
2340 195 200 205  
2342 Pro Glu Pro Ser Thr Ala Pro Ser Thr Ser Phe Leu Leu Pro Met Gly  
2343 210 215 220  
2345 Pro Ser Pro Pro Ala Glu Gly Ser Thr Gly Asp  
E--> 2346 225 230 235

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/10/575,261

DATE: 04/24/2006  
TIME: 16:15:43

Input Set : A:\seq list.txt  
Output Set: N:\CRF4\04242006\J575261.raw

**COPY**

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:17; Line(s) 832  
Seq#:74; Line(s) 2501  
Seq#:75; Line(s) 2558  
Seq#:76; Line(s) 2666

VERIFICATION SUMMARY  
PATENT APPLICATION: US/10/575,261

DATE: 04/24/2006  
TIME: 16:15:43

Input Set : A:\seq list.txt  
Output Set: N:\CRF4\04242006\J575261.raw

COPY

L:9 M:270 C: Current Application Number differs, Replaced Current Application No  
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:48 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:52 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:56 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:60 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:64 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:68 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:72 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:76 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:80 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:84 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:88 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:92 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:97 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:101 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:105 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:109 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:113 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:117 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:121 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:153 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:2  
M:332 Repeated in SeqNo=2  
L:208 M:252 E: No. of Seq. differs, <211> LENGTH:Input:376 Found:372 SEQ:2  
L:584 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7  
L:700 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:8  
L:1396 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:46  
L:1529 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:48  
L:1676 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:51  
L:1786 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:53  
L:2346 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:64